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| 10/076,964 | 02/15/2002 | Herbert F. Cattell | 10010327-1 | 1474 |
| | 7590 07/09/2007 CHNOLOGIES, INC. | | EXAM | INER |
| Legal Department, DL429 Intellectual Property Administration P.O. Box 7599 Loveland, CO 80537-0599 | | | BASOM, BLAINE T | |
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Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

| | Application No. | Applicant(s) | | |
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| | 10/076,964 | CATTELL ET AL. | | |
| Office Action Summary | Examiner | Art Unit | | |
| | Blaine Basom | 2173 | | |
| The MAILING DATE of this communication app Period for Reply | ears on the cover sheet with the o | orrespondence address | | |
| A SHORTENED STATUTORY PERIOD FOR REPLY WHICHEVER IS LONGER, FROM THE MAILING DA - Extensions of time may be available under the provisions of 37 CFR 1.13 after SIX (6) MONTHS from the mailing date of this communication. - If NO period for reply is specified above, the maximum statutory period w - Failure to reply within the set or extended period for reply will, by statute, Any reply received by the Office later than three months after the mailing earned patent term adjustment. See 37 CFR 1.704(b). | ATE OF THIS COMMUNICATION 36(a). In no event, however, may a reply be tin vill apply and will expire SIX (6) MONTHS from 1, cause the application to become ABANDONE | N. nely filed the mailing date of this communication. ED (35 U.S.C. § 133). | | |
| Status | | | | |
| Responsive to communication(s) filed on <u>20 Ju</u> This action is FINAL . 2b)⊠ This Since this application is in condition for allowar closed in accordance with the practice under E | action is non-final. nce except for formal matters, pro | | | |
| Disposition of Claims | | | | |
| 4) ⊠ Claim(s) <u>15-37</u> is/are pending in the application 4a) Of the above claim(s) is/are withdray 5) □ Claim(s) is/are allowed. 6) ⊠ Claim(s) <u>15-37</u> is/are rejected. 7) □ Claim(s) is/are objected to. 8) □ Claim(s) are subject to restriction and/or | vn from consideration. | | | |
| Application Papers | | | | |
| 9) The specification is objected to by the Examine 10) The drawing(s) filed on is/are: a) acce Applicant may not request that any objection to the Replacement drawing sheet(s) including the correct 11) The oath or declaration is objected to by the Ex | epted or b) objected to by the drawing(s) be held in abeyance. Se ion is required if the drawing(s) is ob | e 37 CFR 1.85(a). njected to. See 37 CFR 1.121(d). | | |
| Priority under 35 U.S.C. § 119 | | | | |
| 12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f). a) All b) Some * c) None of: 1. Certified copies of the priority documents have been received. 2. Certified copies of the priority documents have been received in Application No 3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)). * See the attached detailed Office action for a list of the certified copies not received. | | | | |
| Attachment(s) 1) Notice of References Cited (PTO-892) 2) Notice of Draftsperson's Patent Drawing Review (PTO-948) 3) Information Disclosure Statement(s) (PTO/SB/08) Paper No(s)/Mail Date | 4) Interview Summary Paper No(s)/Mail D 5) Notice of Informal F 6) Other: | ate | | |

DETAILED ACTION

Response to Arguments

The Applicants' submission of the Declaration under 37 CFR 1.132, with respect to the rejection of the pending claims has been fully considered. Therefore, the rejection has been withdrawn. However, upon further consideration, a new grounds of rejection is made, as is presented *infra*. Prosecution is hereby reopened.

Claim Rejections - 35 USC § 102

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless -

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

Claims 15, 18, 24-33, and 37 are rejected under 35 U.S.C. 102(b) as being anticipated by the "ScanAlyze" program, as is described by the "ScanAlyze User Manual." In general, ScanAlyze is a program for the analysis of DNA microarrays (see page 4).

Specifically regarding claims 15 and 27, ScanAlyze involves displaying an image of a molecular array, whereby the user can "flag" spots according to various parameters, the flagged spots being indicated by a bold outline over their corresponding location on the array image (for example, see pages 19 and 25). Such flagging is used to indicate one or more characteristics ("Ch1D," "Ch2D," "Ch1B," "Ch2B," "Ch1D/B," "Ch2D/B," "Ch1GTB1," "Ch2GTB1," "Ch1KSD," and/or "Ch2KSD") of the array data (for example, see pages 19 and 25). It is

apparent that the user can arbitrarily choose one or more of these characteristics to flag spots, resulting in a first set of bold outlines being superimposed over the image, and then choose a different one or more of these characteristics, resulting in a second set of bold outlines being superimposed over the array image. As the first set of bold outlines are displayed separately from the second set of bold outlines, the first set of bold outlines are considered graphically distinct from the second set. The user may thus sequentially superimpose two sets of graphically distinct graphical objects (i.e. bold outlines) representing at least two sets of characteristics of the data on the molecular array, wherein graphical objects representing a first set of characteristics are graphically distinct from graphical objects representing a second set of characteristics, and if the user so chooses, each graphically distinct graphical object representing the first set can represent a single characteristic of the data ("Ch1D," "Ch2D," "Ch1B," "Ch2B," "Ch1D/B," "Ch2D/B," "Ch1GTB1," "Ch2GTB1," "Ch1KSD," or "Ch2KSD") that is distinct from the second set of characteristics, each graphically distinct graphical object being superimposed over positions where the data characteristics represented occur on the displayed image of the molecular array (for example, see pages 19 and 25). The positions over which the graphically distinct graphical objects (i.e. bold outlines) are superimposed include feature positions and background positions on the image of the molecular array (see e.g. the figures on pages 4, 18, and 22: the bold outlines are placed over the positions of features, which are themselves located over the background, and therefore, the bold outlines can be considered to be superimposed over both the feature positions and background positions). Accordingly, ScanAlyze is considered to teach a method like that of claim 15. By the same reasoning, the graphical user interface of the ScanAlyze program is considered a graphical user interface like that recited in claim 27.

Regarding claims 18, 28-33, and 37, the characteristics in which the distinct graphical objects are superimposed are selected by the user (see page 19), and are related to the validity of the background, the validity of the feature, and the location of the feature (see page 25). Such graphical objects thus may be used to indicate: a statistically valid feature; a statistically invalid feature; a statistically valid feature background; a statistically invalid feature background; an outlier feature due to non-uniformity of pixel intensities within the feature, due to statistical variance in signal intensity from other features, or due to both non-uniformity of pixel intensities and statistical variance in signal intensities; and outlier feature background due to non-uniformity of pixel intensity with the background, due to statistical variation of the background region from the background regions surrounding other features of the array, or due to both non-uniformity of pixel intensities and statistical various of the background region; or a position of a center of a feature found by analyzing pixel intensities within and near the feature or by row and column indices and a refined feature grid determined from locations of strong features identified. As it is to the user's discretion regarding the use of these distinct graphical objects, it is understood that they may optionally be superimposed only over statistical outlier features and feature backgrounds.

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As per claims 24-26, ScanAlyze teaches reading a sample-exposed array, and visually displaying results using the method described above. It is understood that the user may further process the results, for example, by adjusting parameter values (as done in page 19, for example). As such results are maintained on a computer, presumably via a file, it is understood that such results may be forwarded to a remote location, as is well-known in the art.

Claim Rejections - 35 USC § 103

The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:

(a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negatived by the manner in which the invention was made.

Claims 16-17 and 34-35 are rejected under 35 U.S.C. 103(a) as being unpatentable over the ScanAlyze application, as is described above, over U.S. Patent No. 6,453,251, which is attributed to Bassett, Jr. et al. (hereafter referred to as "Bassett"), and also over U.S. Patent No. 6,437,800, which is attributed to Malamud et al. (and hereafter referred to as "Malamud"). As described above, ScanAlyze teaches a method like that recited in each of claims 15 and 27, whereby a molecular array image is displayed concurrently with feature extraction results associated therewith. It is understood that a user may position a pointer over the position of a feature, to flag a feature for example (for example, see page 18). Regarding the claimed invention, however, ScanAlyze does not explicitly disclose that alphanumeric information related to a selected feature is displayed in response to selecting the feature, as is expressed in claims 16-17 and 34-35. Like ScanAlyze, Bassett describes a program for analyzing microarray data (for example, see column 1, line 36 – column 2, line 10). Bassett particularly teaches displaying an image of a microarray, whereby the user may select a spot within the array, resulting in the display of a panel that presents specific information about the selected spot (for example, see figure 6, and its associated description at column 13, line 66 – column 14, line 13). It would have therefore been obvious to one of ordinary skill in the art, having the teachings of ScanAlyze and Bassett before him at the time the invention was made, to modify the user

interface of ScanAlyze to include functionality of Bassett, so that in response to selecting a spot, alphanumeric data associated with that feature is displayed. It would have been advantageous to one of ordinary skill to utilize this combination because such functionality provides the user with an efficient and easy means for obtaining specified information about any spot, as is demonstrated by Bassett. ScanAlyze and Bassett thus teach receiving an input indication of a feature, i.e. spot, and displaying an alphanumeric representation of information related to the feature, including results from a feature extraction process. Neither ScanAlyze nor Bassett, however, explicitly teach displaying such alphanumeric information within a tooltip, as is claimed. Nevertheless, tooltips are well known in the art. For example, Malamud teaches displaying a tooltip in response to a user positioning a pointer over a graphical object, wherein the tooltip displays alphanumeric information associated with the object (for example, see column 1, lines 34-49; and column 3, lines 26-47). It would have therefore been obvious to one of ordinary skill in the art, having the teachings of ScanAlyze and Malamud before him at the time the invention was made, to modify the user interface of ScanAlyze to include the tooltips of Malamud, so that in response to positioning a cursor over a feature, alphanumeric data associated with that feature is displayed in a tooltip. It would have been advantageous to one of ordinary skill to utilize this combination because such tooltips may reduce confusion and the burden of the user when viewing data associated with a feature, as is taught by Malamud (for example, see column 1, lines 15-43).

Claims 19-23 and 36 are rejected under 35 U.S.C. 103(a) as being unpatentable over the ScanAlyze application, as described above, and also over the "Dapple" application, as described

by the article entitled, "Dapple: Improved Techniques for Finding Spots on DNA Microarrays," which is attributed to Buhler et al. (and hereafter referred to as "Buhler"). As described above, ScanAlyze involves displaying distinct graphical objects superimposed over features of a molecular array. The characteristics in which the distinct graphical objects are superimposed are selected by the user (see page 19), and are related to the validity of the background, the validity of the feature, and the location of the feature (see page 25). Such graphical objects thus may be used to indicate: a statistically valid feature; a statistically invalid feature; a statistically valid feature background; a statistically invalid feature background; an outlier feature due to nonuniformity of pixel intensities within the feature, due to statistical variance in signal intensity from other features, or due to both non-uniformity of pixel intensities and statistical variance in signal intensities; and outlier feature background due to non-uniformity of pixel intensity with the background, due to statistical variation of the background region from the background regions surrounding other features of the array, or due to both non-uniformity of pixel intensities and statistical various of the background region; or a position of a center of a feature found by analyzing pixel intensities within and near the feature or by row and column indices and a refined feature grid determined from locations of strong features identified. ScanAlyze, however, does not involve using distinct types of indications, distinct from the others in terms of shape or color, to indicate such characteristics. Like ScanAlyze, Dapple is an application used for displaying and analyzing molecular arrays (see the "Introduction" on page 1). Dapple particularly teaches marking spots using a plurality of distinct graphical objects, to indicate valid features, invalid features, and "intermediate" quality features (see 3.3 on page 5), and is therefore understood to involve distinct types of indications, distinct from the others in terms of shape or

color, to indicate such characteristics. Accordingly it would have been obvious to one of ordinary skill in the art at the time the invention was made to modify the ScanAlyze application by using different types of indications to indicate valid features, valid feature backgrounds, invalid features, invalid feature backgrounds, and positions of features, as taught by Dapple. One would have been motivated to create such a combination because such different types of indications aid the user in analyzing a molecular array, as is demonstrated by Dapple. As such indications are arbitrary, these indications may comprise figures like recited in claims 20 and 21, and have colors like expressed in claims 22 and 23.

Conclusion

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Blaine Basom whose telephone number is (571) 272-4044. The examiner can normally be reached on Monday through Friday, from 8:30 am to 5:30 pm.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, John Cabeca can be reached on (571) 272-4048. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR

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btb 7/5/2007

JOHN CABECA

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